

EXHIBIT 3

TAVTIGIAN DECLARATION
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CLUSTAL 2.0.12 multiple sequence alignment

USP-5837492 SEQ-ID-NO-2
ICR-Wooster BRCA2_prot
USP-5837492 SEQ-ID-NO-2
ICR-Wooster BRCA2_prot

MPIGSKERPTEEEFKTRCNKADLGPISLNWEELSSSEAPPYNSEAPPNEES 50

EHKNNYEPNLKTPQRKESYNQLASTPIIFKEQGLTLPLYQSPVKELDK 100

[~1,400 intervening amino acids in true BRCA2 structure missing from ICR's "BRCA2" structure.]

USP-5837492_SEQ-ID-NO-2
ICR-Wooster_BRCA2 prot

DQKPEELHNFSLNSELHSDIRKNKMDILSYEETDIVKHKILKESVPUVGTG 1500

USP-5837492_SEQ-ID-NO-2
ICR-Wooster_BRCA2_prot

NQLVTFQGQP1550
PERDEKIKEPTLLGEHTASGKKVVKIAKESLDKVKNLFD7
KE-----KTFMLKE

USP-5837492_SEQ-ID-NO-2
ICR-Wooster_BRCA2_prot

QGTSEITSFSHQWAKTLKYREACKDLELACETIEITAAPKCKEMQNSLNN 1600
QGTSEITSFSHQWAKTLKYREACKDLELACETIEITAAPKCKEMQNSLNN 57

[ICR's "BRCA2" protein sequence aligns with Myriad's sequence, with some isolated errors, for roughly 1,000 amino acids]

USP-5837492_SEQ-ID-NO-2
ICR-Wooster_BRCAA2_Prot

KINSKNAESFOEHTEDYFGKESIWTGGKQIADGGWLIPSNDKGAGKEEF 2600
KINSKNAESFOEHTEDYFGKESIWTGGKQIADGGWLIPSNDKGAGKEEF 1057

USP-5837492_SEQ-1D-NO-2

YRALCDTGPVDPKLIISRIWVNHYRWIIWKLAAMECAFPEKFAANRCLSPPE 2650
YDADGTTTAAAT

USP-5837492 SEQ-ID-NO-2
ICR-Wooster BRCA2 PROT

RVLQLKYRYDTEIDRSRRAIKKIMERDDTAAKTLVLCVSDIISLSANI 2700

Comment [b1]: ICR's "BRCA2" protein sequence diverges from true BRCA2 protein here.

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[~700 intervening amino acids in true BRCA2 structure missing from ICR's "BRCA2" structure.]

USP-5837492_SEQ-ID-NO-2
ICR-Wooster_BRCA2_prot

ALLSGSTGEKQFISVSESTRAPSSSEDYLRLKRRCTTSLIKEQESSQAS 3400

USP-5837492_SEQ-ID-NO-2
ICR-Wooster_BRCA2_prot

TEECEKNKQDTITKKYI 3418